

SEQUENCE LISTING

	(1) GE	NERAL INFORMATION:
5	(i)	APPLICANT: Jardieu, Paula M. Presta, Leonard G.
	(ii)	TITLE OF INVENTION: Humanized Anti-CD11a Antibodies
10	(iii)	NUMBER OF SEQUENCES: 71
15	(iv)	CORRESPONDENCE ADDRESS: (A) ADDRESSEE: Genentech, Inc. (B) STREET: 1 DNA Way (C) CITY: South San Francisco (D) STATE: California (E) COUNTRY: USA (F) ZIP: 94080
20	(v)	COMPUTER READABLE FORM: (A) MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk (B) COMPUTER: IBM PC compatible (C) OPERATING SYSTEM: PC-DOS/MS-DOS (D) SOFTWARE: WinPatin (Genentech)
25	(vi)	CURRENT APPLICATION DATA: (A) APPLICATION NUMBER: (B) FILING DATE: 19-SEP-03 (C) CLASSIFICATION:
30	(vii)	PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: 60/031971 (B) FILING DATE: 27-NOV-1996
35	(vii)	PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: 08/974899 (B) FILING DATE: 20-NOV-1997
40		PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: 09/420745 (B) FILING DATE: 20-OCT-1999
45		PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: 09/975798 (B) FILING DATE: 28-FEB-2001
	(viii)	ATTORNEY/AGENT INFORMATION: (A) NAME: Tan, Lee K. (B) REGISTRATION NUMBER: 39,447

	(ix) TELECOMMUNICATION INFORMATION: (A) TELEPHONE: 650/225-4462 (B) TELEFAX: 650/952-9881						
5	(2) INFORMATION FOR SEQ ID NO:1:						
10	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 108 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY: Linear						
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:						
15	Asp Val Gln Ile Thr Gln Ser Pro Ser Tyr Leu Ala Ala Ser Pro 1 5 10 15						
	Gly Glu Thr Ile Ser Ile Asn Cys Arg Ala Ser Lys Thr Ile Ser 20 25 30						
20	Lys Tyr Leu Ala Trp Tyr Gln Glu Lys Pro Gly Lys Thr Asn Lys 35 40 45						
25	Leu Leu Ile Tyr Ser Gly Ser Thr Leu Gln Ser Gly Ile Pro Ser 50 55 60						
25	Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile 65 70 75						
30	Ser Ser Leu Glu Pro Glu Asp Phe Ala Met Tyr Tyr Cys Gln Gln 80 85 90						
	His Asn Glu Tyr Pro Leu Thr Phe Gly Thr Gly Thr Lys Leu Glu 95 100 105						
35	Leu Lys Arg						
	(2) INFORMATION FOR SEQ ID NO:2:						
40	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 108 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY: Linear						
45	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:						
	Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val 1 5 10 15						

(C) REFERENCE/DOCKET NUMBER: P1014R1C1D1C1

	GIy	Asp	Arg	Val	Thr 20	Ile	Thr	Cys	Arg	A1a 25	Ser	Lys	Thr	Ile	Ser 30
5	Lys	Tyr	Leu	Ala	Trp 35	Tyr	Gln	Gln	Lys	Pro 40	Gly	Lys	Ala	Pro	Lys 45
3	Leu	Leu	Ile	Tyr	Ser 50	Gly	Ser	Thr	Leu	Gln 55	Ser	Gly	Val	Pro	Ser 60
10	Arg	Phe	Ser	Gly	Ser 65	Gly	Ser	Gly	Thr	Asp 70	Phe	Thr	Leu	Thr	Ile 75
	Ser	Ser	Leu	Gln	Pro 80	Glu	Asp	Phe	Ala	Thr 85	Tyr	Tyr	Cys	Gln	Gln 90
15	His	Asn	Glu	Tyr	Pro 95	Leu	Thr	Phe	Gly	Gln 100	Gly	Thr	Lys	Val	Glu 105
	Ile	Lys	Arg												
20	(2) I	NFOF	TAMS	ION 1	FOR S	SEQ :	ID NO	0:3:							
25	(i	(1	A) LI	ENGTI	H: 10	ACTEI 08 ar no Ac	nino		ds						
						Line									
	(xi	.) SI	EQUE	NCE I	DESCI	RIPT:	ION:	SEQ	ID I	10:3	:				
30	Asp 1	Ile	Gln	Met	Thr 5	Gln	Ser	Pro	Ser	Ser 10	Leu	Ser	Ala	Ser	Val 15
35	Gly	Asp	Arg	Val	Thr 20	Ile	Thr	Cys	Arg	Ala 25	Ser	Gln	Ser	Ile	Ser 30
	Asn	Tyr	Leu	Ala	Trp 35	Tyr	Gln	Gln	Lys	Pro 40	Gly	Lys	Ala	Pro	Lys 45
40	Leu	Leu	Ile	Tyr	Ala 50	Ala	Ser	Ser	Leu	Glu 55	Ser	Gly	Val	Pro	Ser 60
	Arg	Phe	Ser	Gly	Ser 65	Gly	Ser	Gly :	Thr	Asp 70	Phe	Thr	Leu	Thr	Ile 75
45	Ser	Ser	Leu	Gln		Glu	Asp	Phe	Ala	Thr 85	Tyr	Tyr	Cys	Gln	Gln 90
					80					03					50

Ile Lys Arg

(2) INFORMATION FOR SEQ ID NO:4:

5	(<i>I</i>		H: 12 Amin			ds						
10	(xi) SF	EQUENCE	DESCR	IPTION	: SEQ	ID N	10:4:	:				
	Glu Val 1	Gln Leu	Gln (Gln Pro	o Gly	Ala	Glu 10	Leu	Met	Arg	Pro	Gly 15
15	Ala Ser	Val Lys	Leu 20	Ser Cy	s Lys	Ala	Ser 25	Gly	Tyr	Ser	Phe	Thr 30
20	Gly His	Trp Met	Asn '	Trp Va	l Arg	Gln	Arg 40	Pro	Gly	Gln	Gly	Leu 45
	Glu Trp	Ile Gly	Met 50	Ile Hi	s Pro	Ser	Asp 55	Ser	Glu	Thr	Arg	Leu 60
25	Asn Gln	Lys Phe	Lys 7	Asp Ly	s Ala	Thr	Leu 70	Thr	Val	Asp	Lys	Ser 75
•	Ser Ser	Ser Ala	Tyr 1	Met Gli	n Leu	Ser	Ser 85	Pro	Thr	Ser	Glu	Asp 90
30	Ser Ala	Val Tyr	Tyr 95	Cys Ala	a Arg	Gly	Ile 100	Tyr	Phe	Tyr	Gly	Thr 105
35	Thr Tyr	Phe Asp	Tyr '	Trp Gl	y Gln	Gly	Thr 115	Thr	Leu	Thr	Val	Ser 120
	Ser						•					
40	(2) INFOR	RMATION EQUENCE								•		
	(E	A) LENGT: B) TYPE: C) TOPOL	Amin		o aci	ds						
45	(xi) SE	EQUENCE :	DESCR	IPTION	: SEQ	ID N	IO : 5 :	:				
٠	Glu Val 1	Gln Leu	Val (Glu Se	r Gly	Gly	Gly 10	Leu	Val	Gln	Pro	Gly 15

	Gly	Ser	Leu	Arg	Leu 20	Ser	Cys	Ala	Ala	Ser 25	Gly	Tyr	Ser	Phe	Thr 30
_	Gly	His	Trp	Met	Asn 35	Trp	Val	Arg	Gln	Ala 40	Pro	Gly	Lys	Gly	Leu 45
5	Glu	Trp	Val	Gly	Met 50	Ile	His	Pro	Ser	Asp 55	Ser	Glu	Thr	Arg	Tyr 60
10	Asn	Gln	Lys	Phe	Lys 65	Asp	Arg	Phe	Thr	Ile 70	Ser	Val	Asp	Lys	Ser 75
	Lys	Asn	Thr	Leu	Tyr 80	Leu	Gln	Met	Asn	Ser 85	Leu	Arg	Ala	Glu	Asp 90
15	Thr	Ala	Val	Tyr	Tyr 95	Cys	Ala	Arg	Gly	Ile 100	Tyr	Phe	Tyr	Gly	Thr 105
20	Thr	Tyr	Phe	Asp	Tyr 110	Trp	Gly	Gln	Gly	Thr 115	Leu	Val	Thr	Val	Ser 120
20	Ser														
25		INFO													
	(() ()		ENGTI YPE :		L3 ar 10 Ac	nino cid		is						
30	(x	i) S	EQUEI	NCE I	DESCI	RIPTI	ON:	SEO	א כדד	TO - C					
									יו עד	NO:6	•				
35	GIu 1	Val	Gln	Leu	Val 5	Glu	Ser					Val	Gln	Pro	Gly 15
35	1				5			Gly	Gly	Gly 10	Leu		Gln Thr		15
35 40	1 Gly	Sér	Leu	Arg	5 Leu 20	Ser	Cys	Gly Ala	Gly Ala	Gly 10 Ser 25	Leu	Phe		Phe	15 Ser 30
	1 Gly Ser	Sér Tyr	Leu Ala	Arg Met	Leu 20 Ser 35	Ser Trp	Cys Val	Gly Ala Arg	Gly Ala Gln	Gly 10 Ser 25 Ala 40	Leu Gly Pro	Phe Gly	Thr	Phe Gly	15 Ser 30 Leu 45
	Gly Ser Glu	Ser Tyr Trp	Leu Ala Val	Arg Met Ser	5 Leu 20 Ser 35 Val 50	Ser Trp Ile	Cys Val Ser	Gly Ala Arg Gly	Gly Ala Gln Asp	Gly 10 Ser 25 Ala 40 Gly 55	Leu Gly Pro Gly	Phe Gly Ser	Thr Lys	Phe Gly Tyr	15 Ser 30 Leu 45 Tyr 60

	Thr A	Ala	Val	Tyr	Tyr 95	Cys	Ala	Arg	Gly	Phe 100	Asp	Tyr	Trp	Gly	Gln 105
5	Gly T	Thr	Leu	Val	Thr 110	Val	Ser	Ser							
Ü	(2) IN	IFOR	MATI	ON E	FOR S	SEQ]	D NO	0:7:							
10	(i)) LE	ENGTH	CHARA I: 18 Amir OGY:	34 an	nino cid	CS: ació	ls						
	(xi)	SE	QUEN	ICE I	DESCF	RIPT	ON:	SEQ	ID 1	10:7	:				
15	Lys G	∃ly	Asn	Val	Asp 5	Leu	Val	Phe	Leu	Phe 10	Asp	Gly	Ser	Met	Ser 15
20	Leu G	3ln	Pro	Asp	Glu 20	Phe	Gln	Lys	Ile	Leu 25	Asp	Phe	Met	Lys	Asp 30
	Val M	1et	Lys	Lys	Leu 35	Ser	Asn	Thr	Ser	Tyr 40	Gln	Phe	Ala	Ala	Val 45
25	Gln F	Phe	Ser	Thr	Ser 50	Tyr	Lys	Thr	Glu	Phe 55	Asp	Phe	Ser	Asp	Tyr 60
·	Val I	ys	Arg	Lys	Asp 65	Pro	Asp	Ala	Leu	Leu 70	Lys	His	Val	Lys	His 75
30	Met I	Leu	Leu	Leu	Thr 80	Asn	Thr	Phe	Gly	Ala 85	Ile	Asn	Tyr	Val	Ala 90
35	Thr G	Glu	Val	Phe	Arg 95	Glu	Glu	Leu	Gly	Ala 100	Arg	Pro	Asp	Ala	Thr 105
	Lys V	/al	Leu	Ile	Ile 110	Ile	Thr	Asp	Gly	Glu 115	Ala	Thr	Asp	Ser	Gly 120
40	Asn I	Ile	Asp	Ala	Ala 125	Lys	Asp	Ile	Ile	Arg 130	_	Ile	Ile	Gly	Ile 135
	Gly I	ùуs	His	Phe	Gln 140	Thr	Lys	Glu	Ser	Gln 145	Glu	Thr	Leu	His	Lys 150
45	Phe A	Ala	Ser	Lys	Pro 155	Ala	Ser	Glu	Phe	Val 160	Lys	Ile	Leu	Asp	Thr 165
	Phe G	Glu	Lys	Leu	Lys 170	Asp	Leu	Phe	Thr	Glu 175	Leu	Gln	Lys	Lys	Ile 180

Tyr Val Ile Glu

(2) INFORMATION FOR SEQ ID NO:8: 5 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 184 amino acids (B) TYPE: Amino Acid (D) TOPOLOGY: Linear 10 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8: Lys Gly Asn Val Asp Leu Ile Phe Leu Phe Asp Gly Ser Met Ser 15 Leu Gln Pro Asp Glu Phe Gln Lys Ile Leu Asp Phe Met Lys Asp 20 Val Met Lys Lys Leu Ser Asn Thr Ser Tyr Gln Phe Ala Ala Val 20 Gln Phe Ser Thr Ser Tyr Lys Thr Glu Phe Asp Phe Ser Asp Tyr Val Lys Gln Lys Asp Pro Asp Ala Leu Leu Glu His Val Lys His 25 Met Leu Leu Thr Asn Thr Phe Gly Ala Ile Asn Tyr Val Ala 30 Thr Glu Val Phe Arg Glu Glu Leu Gly Ala Arg Pro Asp Ala Thr Lys Val Leu Ile Ile Ile Thr Asp Gly Glu Ala Thr Asp Ser Gly 110 120 35 Asn Ile Asp Ala Ala Lys Asp Ile Ile Arg Tyr Ile Ile Gly Ile Gly Lys His Phe Gln Thr Lys Glu Ser Gln Glu Thr Leu His Lys 40 145 Phe Ala Ser Lys Pro Ala Ser Glu Phe Val Lys Ile Leu Asp Thr 155 165 45 Phe Glu Lys Leu Lys Asp Leu Phe Thr Glu Leu Gln Lys Lys Ile

170

Tyr Ala Ile Glu

175

180

	(2) INFORMATION FOR SEQ ID NO:9:
5	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 7 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY: Linear
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:
10	Lys His Val Lys His Met Leu 1 5
	(2) INFORMATION FOR SEQ ID NO:10:
15	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 10 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY: Linear
20	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:
20	Gly Tyr Ser Phe Thr Gly His Trp Met Asn 1 5 10
25	(2) INFORMATION FOR SEQ ID NO:11:
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY: Linear
30	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:
35	Met Ile His Pro Ser Asp Ser Glu Thr Arg Tyr Asn Gln Lys Phe 1 5 10 15 Lys Asp
	пув кор
40	(2) INFORMATION FOR SEQ ID NO:12:
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 amino acids (B) TYPE: Amino Acid (D) TOPOLOGY Linear
45	(D) TOPOLOGY: Linear (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:
	Gly Ile Tyr Phe Tyr Gly Thr Thr Tyr Phe Asp Tyr 1 5 10

5	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 11 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY: Linear
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:
10	Arg Ala Ser Lys Thr Ile Ser Lys Tyr Leu Ala 1 5 10
	(2) INFORMATION FOR SEQ ID NO:14:
15	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 7 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY: Linear
20	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:
	Ser Gly Ser Thr Leu Gln Ser 1 5
25	(2) INFORMATION FOR SEQ ID NO:15:
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 9 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY: Linear
30	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:
	Gln Gln His Asn Glu Tyr Pro Leu Thr 1 5
35	(2) INFORMATION FOR SEQ ID NO:16:
40	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 11 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY: Linear
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:
45	Pro Lys Asn Ser Ser Met Ile Ser Asn Thr Pro

(2) INFORMATION FOR SEQ ID NO:13:

	(2) INFORMATION FOR SEQ ID NO:17:
5	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 7 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY: Linear
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:
10	His Gln Ser Leu Gly Thr Gln 1 5
	(2) INFORMATION FOR SEQ ID NO:18:
15	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 8 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY: Linear
20	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:
	His Gln Asn Leu Ser Asp Gly Lys 1 5
25	(2) INFORMATION FOR SEQ ID NO:19:
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 8 amino acids(B) TYPE: Amino Acid
30	(D) TOPOLOGY: Linear
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:
35	His Gln Asn Ile Ser Asp Gly Lys 1 5
	(2) INFORMATION FOR SEQ ID NO:20:
40	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 8 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY: Linear
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:
45	Val Ile Ser Ser His Leu Gly Gln

	(-,
5	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 26 base pairs(B) TYPE: Nucleic Acid(C) STRANDEDNESS: Single(D) TOPOLOGY: Linear
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:
10	CACTTTGGAT ACCGCGTCCT GCAGGT 26
	(2) INFORMATION FOR SEQ ID NO:22:
15	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 26 base pairs(B) TYPE: Nucleic Acid
	(C) STRANDEDNESS: Single (D) TOPOLOGY: Linear
20	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:
25	CATCCTGCAG GTCTGCCTTC AGGTCA 26
20	(2) INFORMATION FOR SEQ ID NO:23:
30	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY: Linear
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:
35	Met Ile Ala Pro Ala Ser Ser Ser Thr Arg Tyr Asn Gln Lys Phe 1 5 10 15
	Lys Asp
40	(2) INFORMATION FOR SEQ ID NO:24:
45	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 121 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY: Linear

(2) INFORMATION FOR SEQ ID NO:21:

	(xi) SEQUE	NCE DESC	RIPTION	SEQ	ID 1	10:24	1:	:			
	Glu Val Gln 1	Leu Val 5	Glu Sei	Gly	Gly	Gly 10	Leu	Val	Gln	Pro	Gly 15
5	Gly Ser Leu	Arg Leu 20	Ser Cys	: Ala	Ala	Ser 25	Gly	Tyr	Ser	Phe	Thr 30
10	Gly His Trp	Met Asn 35	Trp Val	Arg	Gln	Ala 40	Pro	Gly	Lys	Gly	Leu 45
10	Glu Trp Val	Gly Met 50	Ile Ala	a Pro	Ala	Ser 55	Ser	Ser	Thr	Arg	Tyr 60
15	Asn Gln Lys	Phe Lys 65	Asp Arg	g Phe	Thr	Ile 70	Ser	Val	Asp	Lys	Ser 75
	Lys Asn Thr	Leu Tyr 80	Leu Glr	n Met	Asn	Ser 85	Leu	Arg	Ala	Glu	Asp 90
20	Thr Ala Val	Tyr Tyr 95	Cys Ala	a Arg	Gly	Ile 100	Tyr	Phe	Tyŗ	Gly	Thr 105
25	Thr Tyr Phe	Asp Tyr 110	Trp Gly	/ Gln	Gly	Thr 115	Leu	Val	Thr	Val	Ser 120
23	Ser										
30		NCE CHAR	ACTERIS:	rics:							
	(B) T	ENGTH: 1 YPE: Ami OPOLOGY:		acid	s						
35	(xi) SEQUE	NCE DESC	RIPTION	: SEQ	ID I	NO:2!	5:				
40	Gly Tyr Ser 1	Phe Thr 5	Gly His	Trp	Met	Asn 10					
40	(2) INFORMAT	ION FOR	SEQ ID 1	10:26	:						
45	(B) T	NCE CHAR ENGTH: 1 YPE: Ami: OPOLOGY:	0 amino		s						

	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:
	Gly Tyr Ala Phe Ala Gly Ala Trp Met Asn 1 5 10
5	(2) INFORMATION FOR SEQ ID NO:27:
10	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 10 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:27:
15	Gly Tyr Ala Phe Thr Gly His Trp Met Asn 1 5 10
	(2) INFORMATION FOR SEQ ID NO:28:
20	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 10 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
25	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:28:
	Gly Tyr Ser Phe Ala Gly His Trp Met Asn 1 5 10
30	(2) INFORMATION FOR SEQ ID NO:29:
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 10 amino acids(B) TYPE: Amino Acid
35	(D) TOPOLOGY: (xi) SEQUENCE DESCRIPTION: SEQ ID NO:29:
	Gly Tyr Ser Phe Thr Ala His Trp Met Asn
40	1. 5 10
	(2) INFORMATION FOR SEQ ID NO:30:
45	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 10 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:

	Gly Tyr Ser Phe Thr Gly Ala Trp Met Asn 1 5 10
5	(2) INFORMATION FOR SEQ ID NO:31:
10	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 10 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:31:
15	Gly Tyr Ser Phe Thr Gly His Ala Met Asn 1 5 10
	(2) INFORMATION FOR SEQ ID NO:32:
20	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
25	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:32:
	Met Ile His Pro Ser Asp Ser Glu Thr Arg Tyr Asn Gln Lys Phe 1 5 10 . 15
30	Lys Asp .
	(2) INFORMATION FOR SEQ ID NO:33:
35	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
40	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:33:
	Met Ile Ala Pro Ala Asp Ala Glu Thr Arg Tyr Asn Gln Lys Phe 1 5 10 15
45	Lys Asp

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:

	(2) INFORMATION FOR BEG ID NO.54.
5	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:34:
10	Met Ile His Pro Ser Ala Ser Ala Thr Ala Tyr Asn Gln Lys Phe 1 5 10 15
	Lys Asp
15	(2) INFORMATION FOR SEQ ID NO:35:
20	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:35:
25	Met Ile Ala Pro Ala Asp Ser Glu Thr Arg Tyr Asn Gln Lys Phe 1 5 10 15
	Lys Asp
30	(2) INFORMATION FOR SEQ ID NO:36:
35	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:36:
40	Met Ile His Pro Ala Asp Ala Glu Thr Arg Tyr Asn Gln Lys Phe 1 5 10 15
	Lys Asp
45	(2) INFORMATION FOR SEQ ID NO:37:
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid

(2) INFORMATION FOR SEQ ID NO:34:

	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:37:
5	Met Ile Ala Pro Ser Asp Ala Glu Thr Arg Tyr Asn Gln Lys Phe 1 10 15
	Lys Asp
10	(2) INFORMATION FOR SEQ ID NO:38:
15	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:38:
20	Met Ile Ala Pro Ser Asp Ser Glu Thr Arg Tyr Asn Gln Lys Phe 1 5 10 . 15
	Lys Asp
25	(2) INFORMATION FOR SEQ ID NO:39:
30	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:39:
35	Met Ile Ser Pro Ser Asp Ser Glu Thr Arg Tyr Asn Gln Lys Phe 1 5 10 15
	Lys Asp
40	(2) INFORMATION FOR SEQ ID NO:40:
45	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:

(D) TOPOLOGY:

5	Met Ile His Pro Ala Asp Ser-Glu Thr Arg Tyr Asn Gln Lys Phe 1 5 10 15
	Lys Asp
10	(2) INFORMATION FOR SEQ ID NO:41:
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
15	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:41:
20	Met Ile His Pro Ser Ala Ser Glu Thr Arg Tyr Asn Gln Lys Phe 1 5 10 15
20	Lys Asp
25	(2) INFORMATION FOR SEQ ID NO:42:
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
30	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:42:
35	Met Ile His Pro Ser Asp Ala Glu Thr Arg Tyr Asn Gln Lys Phe 1 5 10 15
00	Lys Asp
40	(2) INFORMATION FOR SEQ ID NO:43:
45	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
43	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:43:
	Met Ile His Pro Ser Asp Asn Glu Thr Arg Tyr Asn Gln Lys Phe 1 5 10 15

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:40:

Lys Asp

	(2) INFORMATION FOR SEQ ID NO:44:
5	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
10	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:44:
	Met Ile His Pro Ser Asp Gln Glu Thr Arg Tyr Asn Gln Lys Phe 1 5 10 15
15	Lys Asp
	(2) INFORMATION FOR SEQ ID NO:45:
20	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
25	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:45:
	Met Ile His Pro Ser Asp Ser Ala Thr Arg Tyr Asn Gln Lys Phe 1 5 10 15
30	Lys Asp
	(2) INFORMATION FOR SEQ ID NO:46:
35	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
40	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:46:
-	Met Ile His Pro Ser Asp Ser Glu Thr Ala Tyr Asn Gln Lys Phe 1 5 10 15
45	Lys Asp

5	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:47:
10	Met Ile His Pro Ser Asp Ser Glu Thr Arg Tyr Asn Ala Lys Phe 1 5 10 15
	Lys Asp
15	(2) INFORMATION FOR SEQ ID NO:48:
20	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:48:
25	Met Ile His Pro Ser Asp Ser Glu Thr Arg Tyr Asn Gln Ala Phe 1 5 10 15
	Lys Asp
00	
30	(2) INFORMATION FOR SEQ ID NO:49: .
35	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:49:
40	Met Ile His Pro Ser Asp Ser Glu Thr Arg Tyr Asn Gln Lys Phe 1 5 10 15
	Ala Asp
45	(2) INFORMATION FOR SEQ ID NO:50:
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 17 amino acids(B) TYPE: Amino Acid

(2) INFORMATION FOR SEQ ID NO:47:

	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:50:
5	Met Ile His Pro Ser Asp Ser Glu Thr Arg Tyr Asn Gln Lys Phe 1 5 10 15
	Lys Ala
10	(2) INFORMATION FOR SEQ ID NO:51:
15	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 11 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:51:
20	Gly Ile Tyr Phe Tyr Gly Thr Thr Tyr Phe Asp 1 5 10
	(2) INFORMATION FOR SEQ ID NO:52:
25	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 11 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
30	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:52:
	Gly Ile Ala Phe Ala Gly Thr Thr Tyr Phe Asp 1 5 10
35	(2) INFORMATION FOR SEQ ID NO:53:
40	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 11 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
40	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:53:
45	Gly Ile Tyr Phe Tyr Gly Ala Ala Phe Asp 1 5 10
	(2) INFORMATION FOR SEQ ID NO:54:

(D) TOPOLOGY:

(i) SEQUENCE CHARACTERISTICS:

,	(=, =======
5	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:54:
	Gly Ile Ala Phe Tyr Gly Thr Thr Tyr Phe Asp 1 5 10
10	(2) INFORMATION FOR SEQ ID NO:55:
15	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 11 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:55:
20	Gly Ile Tyr Phe Ala Gly Thr Thr Tyr Phe Asp 1 5 10
	(2) INFORMATION FOR SEQ ID NO:56:
25	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 11 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
30	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:56: Gly Ile Tyr Phe Tyr Gly Ala Thr Tyr Phe Asp
	1 5 10
35	(2) INFORMATION FOR SEQ ID NO:57:
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 11 amino acids (B) TYPE: Amino Acid
40	(D) TOPOLOGY:
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:57:
45	Gly Ile Tyr Phe Tyr Gly Thr Ala Tyr Phe Asp 1 5 10
	(2) INFORMATION FOR SEQ ID NO:58:
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 11 amino acids

(A) LENGTH: 11 amino acids

(B) TYPE: Amino Acid

(D) TOPOLOGY:

5	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:58:
	Gly Ile Tyr Phe Tyr Gly Thr Thr Ala Phe Asp 1 5 10
10	(2) INFORMATION FOR SEQ ID NO:59:
15	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 11 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
,	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:59:
20	Arg Ala Ser Lys Thr Ile Ser Lys Tyr Leu Ala 1 5 10
	(2) INFORMATION FOR SEQ ID NO:60:
25	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 11 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
30	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:60:
30	Arg Ala Ser Ala Ala Ile Ala Ala Tyr Leu Ala 1 5 10
35	(2) INFORMATION FOR SEQ ID NO:61:
00	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 7 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
40	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:61:
•	Ser Gly Ser Thr Leu Gln Ser
45	1 5
. •	(2) INFORMATION FOR SEQ ID NO:62:
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 7 amino acids

(B) TYPE: Amino Acid
(D) TOPOLOGY:

5	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:62:
	Ala Gly Ala Ala Leu Gln Ser 1 5
10	(2) INFORMATION FOR SEQ ID NO:63:
15	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 7 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:63:
20	Ala Gly Ser Thr Leu Gln Ser 1 5
	(2) INFORMATION FOR SEQ ID NO:64:
25	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 7 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
20	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:64:
30	Ser Gly Ala Thr Leu Gln Ser 1 5
0.5	(2) INFORMATION FOR SEQ ID NO:65:
35	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 7 amino acids (B) TYPE: Amino Acid (D) TOROLOGY
40	(D) TOPOLOGY: (xi) SEQUENCE DESCRIPTION: SEQ ID NO:65:
45	Ser Gly Ser Ala Leu Gln Ser 1 5
40	(2) INFORMATION FOR SEQ ID NO:66:
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 amino acids

(B) TYPE: Amino Acid
(D) TOPOLOGY:

	(B) TYPE: Amino Acid (D) TOPOLOGY:
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:66:
5	Gln Gln His Asn Glu Tyr Pro Leu Thr 1 5
	(2) INFORMATION FOR SEQ ID NO:67:
10	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 9 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
15	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:67:
	Gln Gln His Ala Ala Ala Pro Leu Thr 1 5
20	(2) INFORMATION FOR SEQ ID NO:68:
25	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 9 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:68:
30	Gln Gln Ala Asn Glu Tyr Pro Leu Thr 1 5
	(2) INFORMATION FOR SEQ ID NO:69:
35	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 9 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
40	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:69:
40	Gln Gln His Ala Glu Tyr Pro Leu Thr 1 5
45	(2) INFORMATION FOR SEQ ID NO:70:
TŲ	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 9 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:

	(XI) SEQUENCE DESCRIPTION: SEQ ID NO:/0:
5	Gln Gln His Asn Ala Tyr Pro Leu Thr 1 5
	(2) INFORMATION FOR SEQ ID NO:71:
10	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 9 amino acids(B) TYPE: Amino Acid(D) TOPOLOGY:
15	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:71
	Gln Gln His Asn Glu Ala Pro Leu Thr 1 5